

Antibodies to *Cryptosporidium*Technical Field

The present invention relates to antibodies to *Cryptosporidium* and methods to raise suitable *Cryptosporidium*-specific antibodies in animals.

5 Background Art

The protozoan parasite *Cryptosporidium* is amongst the most common pathogens responsible for diarrhoeal disease in humans. Infection occurs when *Cryptosporidium* oocysts shed in the faeces of infected individuals are ingested by new hosts. Recently, several large outbreaks of cryptosporidiosis have occurred in which drinking water has been identified as the source of infection. Surveys have shown that many surface water supplies are contaminated with *Cryptosporidium* oocysts.

Laboratory methods used to detect *Cryptosporidium* often involve the use of antibodies to this organism. Typical methods used to analyse water samples for the presence of this organism include microscopy and cytometry or a combination of these techniques. Flow cytometric methods involve staining of samples with a fluorescently labelled monoclonal antibody specific to the surface of *Cryptosporidium* oocysts and then analysis with a sorter flow cytometer. Particles with the fluorescence and light scatter characteristics of *Cryptosporidium* oocysts are sorted onto a microscope slide and examined manually using epifluorescence microscopy to confirm their identity as oocysts. This confirmation step is necessary because with a single antibody the cytometer is unable to distinguish oocysts from all other particles present in water samples. The particles that the cytometer can mistake as oocysts are autofluorescent particles such as algae or particles that non-specifically bind the oocyst-specific antibody.

Analysis-only flow cytometers are available which are simple to operate and relatively inexpensive. These cytometers are unable to perform sorting. To enable the detection of *Cryptosporidium* oocysts using an analysis-only cytometer the discrimination achieved by the cytometer must be improved so that non-oocyst particles are not mistaken as oocysts. The present inventors have shown previously that it is possible to detect a single specific microorganism in turbid water samples with an analysis cytometer if the microorganism is labelled with two different antibodies.

Unfortunately, the antibodies for *Cryptosporidium* presently available are not ideal due their stickiness and there is a need for more specific and

reactive antibodies to the surface of *Cryptosporidium* oocysts. Monoclonal antibodies (mAbs) that are specific to the surface of *Cryptosporidium* oocysts are used for detecting *Cryptosporidium* in clinical and environmental samples. All available mAbs that bind to the surface of *Cryptosporidium* oocysts are of the immunoglobulin M (IgM) or IgG3 subclass. Monoclonal antibodies of the IgG1 subclass would be preferable because they usually show less non-specific binding. Such mAbs would be more suitable in methods currently used for the detection and identification of *Cryptosporidium*. Unfortunately, past attempts by workers in the field to produce IgG1 monoclonal antibodies to *Cryptosporidium* have been unsuccessful or not substantiated (Smith, 1994; MacDonald *et al.*, 1991). It is generally considered that due to the antigenic characteristics of this organism, this class of antibody is not produced by infected or immunised animals (Smith, 1994).

In WO 97/08204 filed by the present inventors, monoclonal antibodies to a range of *Cryptosporidium* oocyst antigens were developed. Whole or excysted oocysts that were exposed to various treatments were used as antigens. From a total of eight fusions that included screening several thousand hybridomas only one hybridoma was identified that was specific to the surface of *Cryptosporidium* oocysts. This monoclonal antibody was of the IgM immunological subclass.

The present inventors have now developed a new method that allows the production of IgG1 antibodies to the surface of *Cryptosporidium* oocysts.

#### Disclosure of Invention

In a first aspect, the present invention consists in a method of producing IgG1 subclass antibodies reactive to the surface of *Cryptosporidium* oocysts comprising:

- (a) pretreating *Cryptosporidium* oocysts with a reagent so as to remove the surface layer of the oocysts to form an oocyst antigen preparation;
- (b) separating the oocysts from the antigen preparation so as to obtain a separated oocyst antigen preparation capable of eliciting a detectable IgG1 immune response in an animal to the surface of the oocyst;
- (c) immunising an animal with the separated oocyst antigen preparation so as to elicit an IgG1 immune response in the animal; and
- (d) obtaining from the animal IgG1 antibodies reactive to the surface of *Cryptosporidium* oocysts.

It will be appreciated that once a suitable immune response has been stimulated in an animal, for example in a laboratory mouse, monoclonal antibodies of IgG1 subclass may be generated by standard techniques from that animal.

5 In a preferred embodiment of the first aspect of the present invention, the reagent used to prepare the antigen preparation is a detergent, preferably the detergent is sodium dodecyl sulphate (SDS). One suitable pretreatment involves boiling the oocysts in the presence of SDS for a sufficient time to generate a suitable antigen preparation. When a concentration of 0.5% (w/v)  
10 SDS is used, boiling for 1 hour has been found to be particularly suitable.

Other suitable reagents include urea, detergents such as Triton X-100 or nonident. enzymes such chitinase, oxidising agents such as sodium hypochlorite, sodium periodate, ozone and reducing agents such as mercaptol ethanol and 1,1,1-trichloro- 2,2-bis[4-chlorophenyl]ethane (DDT).

15 The pretreatment removes antigens from the surface of the oocyst in a form that will allow the generation of IgG1 antibodies when injected into an animal.

The animal may be immunised by any technique suitable for eliciting an immune response in an animal. Adjuvants may also be included with the  
20 antigen preparation prior to immunising the animal to promote a strong immune response in the animal.

In a further preferred embodiment, the antigen preparation also enhances the production of IgM antibodies when placed in an animal.

25 In a second aspect, the present invention consists in a method of producing IgG1 subclass antibodies reactive to the surface of *Cryptosporidium* oocysts, the method comprising:

- (a) separating at least a portion of the *Cryptosporidium* oocyst wall from the internal sporozoites to form an oocyst-wall preparation;
- (b) treating the separated oocyst-wall preparation so as to obtain an oocyst  
30 antigen preparation capable of eliciting a detectable IgG1 immune response in an animal to the surface of the oocyst;
- (c) immunising an animal with the oocyst antigen preparation so as to elicit an IgG1 immune response in the animal; and
- (d) obtaining from the animal IgG1 antibodies reactive to the surface of  
35 *Cryptosporidium* oocysts.

The present inventors have found that in order to obtain a suitable oocyst wall antigen preparation, the oocyst wall should be separated from internal sporozoite components. It appears that the internal sporozoite antigens are more immunodominant than oocyst wall antigens and their presence in an antigen preparation may mask the oocyst wall antigens. A mixed antigen preparation will usually result in raising antibodies to the sporozoite antigens.

The separation of the oocyst wall from the internal sporozoite (step (a)) can be achieved by any means. The present inventors have found that causing the oocyst to excyst followed by immuno-separation of the wall components is particularly suitable. Separation can also be achieved by surface labelling of whole oocysts with a ligand, such as biotin, allowing separation of cell wall fragments from internal components by reaction with a reagent reactive to the ligand, such as avidin, on an insoluble matrix or beads. It will be appreciated, however, that other separation methods known to the art would also be suitable. Examples include centrifugation, flow cytometry, density gradient separation, precipitation, immuno-labelling, ligand-binding, biotin-labelling and separation by avidin, and chromatographic separation.

It is not necessary to cause the oocyst to excyst by normal procedures. The oocysts can be freeze-thawed for example to promote initial separation of the wall from the internal sporozoites. Furthermore, the oocyst may be physically broken up by crushing, sonication, or grinding followed by separation.

The treating step (b) can be carried out by any means suitable. In particular, the present inventors have found that physically breaking up the cell wall can produce a good antigen preparation. This can be done by any means but the use of a bead beater is quite suitable.

The treatment removes antigens from the surface of the oocyst wall in a form that will allow the generation of IgG1 antibodies when injected into an animal.

It will be appreciated that once a suitable immune response has been stimulated in an animal, for example in a laboratory mouse, monoclonal antibodies of IgG1 subclass may be generated by standard techniques from that animal.

The animal may be immunised by any technique suitable for eliciting an immune response in an animal. Adjuvants may also be included with the antigen preparation prior to immunising the animal to promote a strong immune response in the animal.

5 In a further preferred embodiment, the antigen preparation also enhances the production of IgM antibodies when placed in an animal.

As the present inventors have determined methods that allow the production of useful IgG1 antibodies reactive to the surface of *Cryptosporidium* oocysts, it will be appreciated that similar antibodies to those produced by the present inventors may now be produced from the information and teaching provided herein.

10 In a third aspect, the present invention consists in substantially isolated IgG1 antibodies reactive to the surface of *Cryptosporidium* oocysts produced by the method according to the first or second aspects of the present invention.

15 Preferably, the antibodies are monoclonal antibodies.

In a fourth aspect, the present invention consists in a substantially isolated IgG1 antibody reactive to the surface of *Cryptosporidium* oocysts, the antibody having the oocyst binding and affinity characteristics of antibody CRY104.

20 Preferably, the antibody is a monoclonal antibody.

More preferably, the IgG1 monoclonal antibody is produced by clone CRY104.

25 In a fifth aspect, the present invention consists in the hybridoma clone CRY104.

The production of antibody CRY104 is defined in the specification such that persons skilled in the art will be able to produce other antibodies with *Cryptosporidium* oocyst binding and affinity characteristics similar to, or the same as, CRY104. CRY104 is recited in the specification as one example of a suitable IgG1 antibody reactive to the surface of *Cryptosporidium* oocysts that can be obtained from the teaching of the present invention. Accordingly, a sample of CRY104 is not required for a person to carry out the present invention.

35 Throughout this specification, unless the context requires otherwise, the word "comprise", or variations such as "comprises" or "comprising", will be understood to imply the inclusion of a stated element, integer or step, or

group of elements, integers or steps, but not the exclusion of any other element, integer or step, or group of elements, integers or steps.

In order that the present invention may be more clearly understood, preferred forms will be described with reference to the following examples and the accompanying drawing.

#### Brief Description of Drawings

Figure 1 shows the comparison of monoclonal antibodies for staining oocysts in water samples.

Figure 2 shows flow cytometric analysis of *Cryptosporidium* oocysts seeded in a composite water concentrate. The fluorescent oocyst population is clearly separated from autofluorescent and extraneous particles bound non-specifically to antibody. The optimal concentration of mAb to use for water analysis needs to define a clear separation from background fluorescence as seen in this figure.

Figure 3 shows flow cytometric analysis of a control sample of fluorescently stained *Cryptosporidium* oocysts. A default size elliptical region is centred on the main population of pure fluorescent oocysts for each antibody (R1). A polygon region is defined around the bead standard population (R2).

Figure 4 shows flow cytometric analysis of a water concentrate containing no oocysts or cysts. The regions are those from Figure 2 for each mAb examined. The number of particles within the oocysts or cyst region (R1) contain both autofluorescent and non-specifically bound particles. The number of particles in R1 are divided by the number of beads analysed (R2). The result is a ratio of non-specific binding comparable between antibodies.

#### Modes for Carrying Out the Invention

##### MATERIALS AND METHODS

##### *Cryptosporidium* oocysts

*Cryptosporidium parvum* oocysts were purified from pooled faeces of naturally infected neonatal calves in Sydney. Faecal samples were centrifuged (2000 g, 10 min) and resuspended in water twice and then resuspended in 5 volumes of 1% (w/v) NaHCO<sub>3</sub>. Fatty substances were then extracted twice with 1 volume of ether, followed by centrifugation (2000 g for 10 min). Pellets were resuspended in water and filtered through a layer of pre-wetted non-adsorbent cotton wool. The eluate was then overlaid onto 10 volumes of 55% (w/v) sucrose solution and centrifuged (2000 g for 20 min).

Oocysts were collected from the sucrose interface and the sucrose flotation step repeated until no visible contaminating material could be detected. Purified oocysts were surface sterilised with ice cold 70% (v/v) ethanol for 30 min, washed once in phosphate buffered saline (PBS; Oxoid, Sydney) and stored in PBS at 4°C for up to 2 weeks.

### Antigen preparation

#### Surface Extraction:

A 2 ml sample of oocysts containing approximately  $2 \times 10^9$  oocysts was centrifuged at 13000 g for 1 minute and the supernatant removed and discarded. Oocysts were resuspended in 2 ml of ice cold 0.5% (w/v) SDS and placed in a boiling water bath for one hour. The sample was then centrifuged at 13000 g for 20 minutes to remove the oocysts. The supernatant was carefully removed, mixed with 10 ml of acetone and placed at -20°C for 8 hours. The sample was then centrifuged at 13000 g for 10 minutes and the supernatant discarded. A small white precipitate was then resuspended in sterile PBS.

The protein concentration was measured using the commercially available Biorad DC protein assay using the standard protocol and bovine serum albumin (BSA) as a standard.

#### Oocyst Walls:

*Cryptosporidium* oocysts were excysted as described by Robertson *et al.* (1993). A 1 ml sample of oocysts containing approximately  $1 \times 10^9$  oocysts was centrifuged at 13000 g for 1 minute and the supernatant removed and discarded. Oocysts were resuspended in 1 ml of acidified Hank's balanced salt solution (HBSS), pH 2.7, and incubated at 37°C for 30 min. Samples were then washed and resuspended in 1 ml of HBSS with 100 µl of 1% (w/v) sodium deoxycholate in Hank's minimal essential medium (HMEM) and 100 µl of 2.2% (w/v) NaHCO<sub>3</sub> in HBSS, and incubated at 37°C for 4 h. The sample was then analysed using the Coulter Elite flow cytometer as described previously (Vesey *et al.*, 1997). The population with the lowest forward angle light scatter signal was sorted into test tubes and concentrated by centrifuging at 3000 g for 20 minutes. Concentrated samples were stored at -20°C in PBS.

#### Excystation of Oocysts:

Excystation was achieved as described above for oocysts walls.

#### Purification of Oocyst Walls:

The oocysts walls were purified from the excysted sample using immuno-magnetic separation. A 0.5 ml aliquot of magnet beads (approximately  $5 \times 10^7$  beads) coated with a goat anti-mouse IgM antibody (DYNAL Pty Ltd, Australia) were mixed with 10 ml of tissue culture supernatant of a *Cryptosporidium* oocyst-specific monoclonal antibody CRY26 (Vesey 1996). The beads were incubated at 4°C for 4 hours and then placed next to a magnet so that the beads were drawn to the bottom of the tube. The supernatant was removed and discarded and the beads resuspended in 10 ml of PBS plus 2% (w/v) bovine serum albumin (BSA; Sigma fraction V)(PBS-BSA). This washing procedure was repeated twice and the beads resuspended in a final volume of 1 ml of PBS-BSA. The beads were then mixed with the sample of excysted oocysts and incubated on a rotary shaker at room temperature for 30 minutes. The tube was placed next to the magnet so that the beads and the attached oocysts were attracted to the bottom of the tube. The supernatant was removed and placed at 4°C. To remove any contaminating sporozoites the beads were gently resuspended in 1 ml of PBS-BSA and then concentrated once more using the magnet. The supernatant was removed and discarded. The beads were resuspended in 1 ml of PBS and vortexed vigorously to unattach the beads from the oocyst walls. The beads were concentrated using the magnet and the supernatant containing the oocyst walls removed and kept on ice. The beads were then added to the original sample of excysted oocysts and the entire procedure repeated 10 times. The 10 samples of purified oocyst walls were then pooled and concentrated by centrifuging at 3000 g for 10 minutes. A fraction of the sample of oocyst walls was analysed using flow cytometry as described previously (Vesey et al., 1997). The sample was analysed in a tube containing an exact number of beads (TrueCount, Becton Dickinson, San Hose, USA) to determine the number of oocyst walls.

#### Breaking up of Oocysts Walls:

Half the sample of oocyst walls were treated to break the walls into small pieces using a FastPrep bead beater (Bio101, CA, USA) fifteen times at maximum speed for 40 second durations. The sample was cooled on ice for 1 minute between each 40 second treatment. The oocyst wall pieces were resuspended into 3 ml of PBS and aliquoted into 200 µl amounts and stored frozen until used.



### Immunisation of mice

#### Method 1:

Five Balb/C female mice were immunised by IP injection with either 200 µg of the oocyst surface extract (group E mice) or  $4 \times 10^4$  oocyst walls (group W mice). Antigen preparations were emulsified in Freund's complete adjuvant. Mice were bled prior to receiving the primary injection to provide a negative control. Two further IP injections with the same amount of antigen but emulsified in Freund's incomplete adjuvant were given at 3 week intervals. Mice were bled after the second of these injections to check for immune response. Group E mice were given two final intravenous boosts of 200 µg of antigen were given 3 days and 1 day prior to the fusion.

#### Method 2:

Five Balb/C female mice (group WP) were immunised by IP injection with 200 µl of smashed oocyst wall preparation emulsified in Freund's complete adjuvant. The preparation contained approximately  $1 \times 10^6$  oocyst walls. A second group of mice (group WI) were immunised with approximately  $1 \times 10^6$  intact purified oocyst walls emulsified in Freund's complete adjuvant. Mice were bled by tail bleeding prior to receiving the primary injection to provide a negative control. Two further IP injections with the same amount of antigen but emulsified in Freund's incomplete adjuvant were given at 3 week intervals. Mice were bled after the second of these injections to check for immune response.

The mouse may be given either IP or IV booster injections to 7 days prior to sacrifice and fusion of spleen cells to assist in the development of appropriate antibodies.

#### Analysis of mouse serum

Samples (approximately 50 µl) of blood was collected by tail bleeding and then centrifuged at 13000 g for 30 seconds. The top layer of serum carefully removed and stored at -20°C until analysed. Serum was diluted to 1 in 100, 1 in 1000 and 1 in 10,000, in 1% (w/v) bovine serum albumin in PBS (BSA-PBS). Aliquots (50 µl) of diluted serum were mixed with 10 µl of oocyst suspension in PBS (containing approximately  $1 \times 10^6$  oocysts) and incubated at room temperature for 20 minutes. Samples were mixed with either 50 µl of goat anti-mouse IgM specific antibody conjugated with FITC (diluted 1 in 50 with BSA-PBS)(Sigma product number) or with 50 µl of goat anti-mouse IgG specific antibody conjugated with PE (diluted 1 in 200 with

BSA-PBS)(Sigma). After a further 20 minutes incubation at room temperature samples were analysed using a FACScan flow cytometer. A negative control of PBS and an IgM positive control of tissue culture supernatant from a IgM monoclonal antibody specific to the surface of *Cryptosporidium* oocysts were analysed with each batch of samples. The mean fluorescence intensities of the FITC and the PE stained samples were recorded.

Samples of mouse serum diluted 1 in 500 in BSA-PBS were analysed using western blotting.

#### Production of hybridomas

Mice were sacrificed, spleen cells dissected and fused with NS1 mouse myeloma cells and the resulting hybridomas cloned. A 50 µl volume of tissue culture supernatant from each hybridoma was mixed with 10 µl of oocyst suspension in PBS (containing approximately  $1 \times 10^6$  oocysts).

Samples were incubated at room temperature for 20 minutes and then mixed with 50 µl of goat anti-mouse antibody specific to both IgM and IgG antibodies and conjugated with FITC (diluted 1 in 100 with BSA-PBS)(Silenus, Melbourne). After a further 20 minutes incubation at room temperature samples were analysed using a FACScan flow cytometer. A negative control of tissue culture supernatant from a *Dictyostelium*-specific antibody (MUD62) and a positive control of tissue culture supernatant from a IgM monoclonal antibody specific to the surface of *Cryptosporidium* oocysts (CRY26) were analysed with each batch of samples. Hybridomas that produced a higher mean fluorescence (FL1) than the negative control were cloned and tested once more.

The immunological subclass of monoclonal antibody produced by clones was identified using the Sigma Immuno Type Kit.

#### Hybridoma Screening

Approximately 7-14 days after the fusion microwell plates were monitored for hybridoma growth. Hybridomas visible at 40x objective were marked, labelled and 100µl of tissue culture supernatant aseptically removed without disturbing the hybridomas at the bottom. All hybridomas were then refed with 100µl of fresh medium for continued growth.

#### Initial Screen:

To each 100 µl of hybridoma supernatant collected 10 µl of  $1 \times 10^8$  oocysts was added and allowed to incubate for 15 minutes at room

temperature. A second antibody anti-mouse FITC (Amrad) was then added (100 µl at 1:50 dilution) and incubated at room temperature for 15 minutes. Samples were then analysed by flow cytometry. Fluorescence intensity of the oocyst population was measured. High fluorescence seen on histogram indicated a positive for anti-*Cryptosporidium* antibodies. All positives were then marked and grown up for further culturing.

#### Secondary Screen:

Once positive hybridomas had been culture into 12 well microtitre plates they were then tested for IgM or IgG antibody class. A 100 µl sample of each positive was placed in two separate tubes. To each tube 10 µl of  $1 \times 10^8$  oocysts was added and incubated for 15 minutes at room temperature. To each duplicate tube 100 µl of prediluted FITC labelled, anti-IgG (Zymed 61-6011), or FITC labelled anti-IgM (Sigma F-9259) was added and allowed to incubate at room temperature for 15 minutes. Each sample was then analysed by flow cytometry. High fluorescence of the oocyst population observed on a histogram, indicated a positive for the antibody subclass which would later be confirmed.

#### Subclass Confirmation:

All positive hybridomas were confirmed for Isotype by a commercially available (Serotec) haemagglutination assay, which employs sheep erythrocytes conjugated with an antibody which specifically recognises a mouse Ig isotype.

#### Analysis of antibodies for oocyst staining in water samples

The effectiveness of mAbs for use in water samples was evaluated by flow cytometry. A 100 µl volume of supernatant was added to seeded samples. Seeded samples consisted of 50 µl water concentrate and 10µl of a high oocyst seed. This was allowed to incubate at room temperature for 20 minutes. After this incubation 100 µl volume of anti-mouse FITC conjugated antibody (Silenus, 1:50) was added and incubated for a further 20 minutes. Samples were then analysed using flow cytometry. Data was analysed to determine which antibody produced the greatest separation between the immunofluorescent control (oocyst) population and the background fluorescent particles detected within water concentrates.

#### Functional Measurement of Avidity

Avidity of binding (affinity constant of whole antibody) of the FITC labelled anti-*Cryptosporidium* antibodies CRY104, CRY26 and a commercial

anti-*Cryptosporidium* FITC-labelled antibody (Immunocel) were measured as follows. Antibodies of a known concentration were serially diluted out from a 20 µg concentration down for 20 two-fold dilutions. Each concentration of mAb was then incubated with  $1 \times 10^7$  oocysts for 20 minutes at room temperature. A negative control of oocysts in PBS was also prepared to provide an end point for binding. Fluorescence (FL-1) values for each dilution were recorded and plotted. The value for 50% maximal binding to the oocysts for each mAb analysed was obtained. Assumptions were made that the total input antibody is very nearly the same as free antibody, therefore the dissociation constant (kd) is equal to this (50%) concentration. The affinity constant (ka) is then calculated as the reciprocal value.

#### Flow cytometry

A FACScan flow cytometer was used for analysis of mouse serum and hybridomas. Logarithmic signals were used for all detectors. The threshold was set on side scatter at a value of 500. The detectors were set at the following levels of sensitivity: 200 for side scatter (SSC); E00 for forward scatter (FALS); 600 for the green fluorescence detector (FL1) and 600 for the red fluorescence detector (FL2). A region (R1) was defined on a dot plot of FALS versus SSC that enclosed single oocysts but not clumps of oocysts. Histograms of FL1 and FL2 were gated so that the only particles that appeared in region R1 would appear on the histograms. The mean value of FL1 and or FL2 from the histograms were recorded for 2000 oocysts from each sample analysed.

#### Sodium Dodecyl sulphate-polyacrylamide electrophoresis (PAGE)

Two hundred microlitres of oocysts at  $5 \times 10^7$  -  $10^8$  cells/ml were added to an equal volume of reducing buffer (composed of 975 µl 0.125 M Tris HCl/0.5% (w/v) SDS, 150 µl glycerol, 225 µl 10% (w/v) SDS, 150 µl 2-Mercaptoethanol and 20 µl Bromophenol Blue (0.25% w/v) and boiled for 3 minutes at 100°C.

This reduced sample was then run on a 12% polyacrylamide separating gel with a 5% stacking gel in a Biorad Miniprotean II cell apparatus. Proteins for high and low molecular weight markers (Novex) were run alongside the sample for approximately 45 - 60 minutes at 200 volts.

#### Western Blotting

Aliquots (100 µl) of oocyst suspension in PBS containing approximately  $5 \times 10^7$  oocysts were mixed with 100 µl of reducing buffer and

boiled for 2 minutes. The entire sample was loaded onto a 10% SDS slab gel and run at 150 V for 1 hour. A sample of prestained molecular weight markers (Novex) was included. Following SDS-PAGE, the proteins were transferred to nitrocellulose using a semi-dry electroblotting system and a discontinuous buffer system. The nitrocellulose was then cut into 2 mm wide strips and soaked in 2% (w/v) skimmed milk in PBS. The strips were then incubated with either mouse serum or hybridoma culture supernatant. Tissue culture supernatant from an IgM *Cryptosporidium*-specific monoclonal antibody was included as a positive control. Samples were incubated for 1 hour at room temperature. The nitrocellulose strips were then rinsed for 3 x 5 min in 3% (w/v) skimmed milk powder in PBS; then incubated for 1 h in a goat anti-mouse antibody (specific to both IgM and IgG antibodies) conjugated to horse radish peroxidase (HRP, Tago, Inc. Burlingame, California, USA), diluted to 1:1500 with 3% (w/v) skimmed milk powder in PBS. Strips were then washed for 3 x 5 min in PBS and developed using a fresh solution of 4CN substrate and finally washed extensively under running tap water for at least 20 min.

#### Water samples

Samples (10 L) of untreated surface water were collected from locations around Australia and concentrated using a flocculation technique (Vesey *et al.* 1993a). A composite untreated surface water sample was prepared by mixing aliquots of samples from 15 different sites. The sample was centrifuged at 3000 g for 10 min and the pellet resuspended in PBS. Aliquots (50 µl) of water sample concentrate were seeded with 10 µl of oocyst suspension (containing approximately 1000 oocysts) and mixed thoroughly. The samples were then mixed with 50 µl of tissue culture supernatant from a *Cryptosporidium*-specific monoclonal antibody and incubated for 20 minutes at room temperature. An aliquot (50 µl) of a FITC conjugated goat anti-mouse antibody (specific to both IgG and IgM antibodies)(Silenus) diluted 1 in 300 in PBS-BSA was then added to each sample and incubated for 20 minutes at room temperature. Samples were then analysed using flow cytometry.

#### ALTERNATIVE METHODS

To separate the oocyst walls from the sporozoites the oocysts can be freeze-thawed instead of excysted. A purification step such as immuno-magnetic separation, flow cytometry or density gradient separation is

required to purify the oocyst walls away from the sporozoites. The present inventors have prepared cell wall samples by surface labelling whole oocysts with biotin and after excystation binding the cell walls to avidin-labelled magnetic beads.

5           One alternative approach would involve smashing the whole oocysts (sporozoites still inside) into small pieces and then using an immunological purification method such as immuno-magnetic separation or affinity chromatography.

10           Other alternative methods such as sonication could be used to break up the oocysts or the purified oocyst walls.

#### ANTIBODY COMPARISON TESTS

##### *Cryptosporidium* oocysts and *Giardia* cysts

15           *Cryptosporidium parvum* oocysts were isolated from naturally infected calves as previously described. Oocysts were heat inactivated at 65°C for 15 minutes. *Giardia lamblia* cysts were obtained formalin fixed from Waterborne, Inc. (New Orleans, USA). Cysts and oocysts were stored in phosphate buffered saline (0.01M phosphate, pH 7.3 ± 0.2) (Oxoid) at 4°C.

##### *Cryptosporidium*-specific and *Giardia*-specific antibodies

20           Seven *Cryptosporidium*-specific and four *Giardia*-specific mAbs were evaluated in this study, their supplier, fluorescent label and class of antibody are described in Table 3 and 4.

##### Concentrated water samples

25           Samples (10 L) of untreated surface water were collected from sites around Australia and concentrated by flocculation or filtration. A composite untreated water sample was prepared by mixing aliquots of samples from a range of different sites. The sample was centrifuged at 13,000 rpm for 10 min and the pellet resuspended in PBS. The volume of the sample was adjusted so that 100 µl of concentrate was equivalent to 5 litres of original untreated water. Samples were also prepared from different water types which include  
30           raw river, effluent, filtered and backwash. All samples were pre-filtered through a 38-micron stainless steel mesh filter prior to use.

##### Sample preparation

35           Fluorebrite™ beads (Polysciences, Inc. Warrington, PA, USA.) were used to standardise the assay. The beads were added at a high concentration into bovine serum albumin (BSA, Sigma Chemical Co, St Louis USA) (4% w/v) in PBS and azide (0.05% v/v). The same volume of beads was then

added to every sample, making it possible to monitor the volume of concentrate analysed, by the number of beads detected. The volume of water concentrate analysed for each sample was then standardised by producing a result per bead analysed. Sample buffer for staining samples consisted of

5 BSA (1% w/v) Tween20 (0.05%v/v) in PBS.

#### **Antibody concentrations**

The optimal concentration or dilution of each antibody to be evaluated (Table 3 and 4) was firstly determined by flow cytometry. Serial dilutions for mAbs of unknown protein concentration were setup between 1:20 and

10 1:1,280 into appropriate sample buffer as described by the manufacturer. Those mAbs of known protein concentrations were set-up as serial dilutions between 8 µg/ml to 0.5 µg/ml. A 100 µl aliquot of each dilution was added to seeded samples. Seeded samples consisted of 50 µl water concentrate and 50 µl of a high oocyst or cyst seed. Samples were incubated at room

15 temperature for 30 minutes and then analysed using flow cytometry. Data was analysed to determine which concentration of antibody produced the greatest separation between the immunofluorescent control (oocyst/cysts) population and the background fluorescent particles detected within water concentrates (Figure 2).

#### **Staining water samples**

Following optimisation of antibody concentrations, each mAb was diluted to the appropriate concentration in sample buffer. All samples were prepared in 5 ml Falcon tubes. Positive controls were prepared consisting of 50 µl of the respective high seed. Water samples to be tested were prepared

25 in triplicate and consisted of 50 µl water concentrate. Each mAb was added to controls and water samples to a total volume of 150 µl in sample buffer and mixed. After incubation at room temperature for 30 minutes samples were seeded with 20 µl aliquots of the Fluorbrite™ bead standard. Samples were then mixed by vortexing for 5 seconds and analysed.

30 The Hydrofluor mAb kit contains unconjugated *Cryptosporidium*-specific and *Giardia*-specific antibodies as well as an anti-mouse FITC conjugated antibody. The reagents were used as recommended in the manufacturers instructions.

#### **Flow Cytometry**

35 Comparison of *Cryptosporidium* and *Giardia* mAbs was carried out using a Becton-Dickinson FACScan flow cytometer (Becton-Dickinson, Lane

Cove, NSW Australia). Sheath fluid consisted of undiluted Isoton II (Coulter Electronics, Brook Vale, NSW Australia). Detectors used were side angle light scatter (SSC) versus FL1 detector (green fluorescence). Voltages of detectors were set on 300 mV for SSC and 450 mV for all remaining detectors. Threshold was set on fluorescence detector 1 (FL1). Compensation was set at FL1-FL2 45%. Positive samples for each mAb were analysed first. Approximately 1,000 events were collected, and an ellipse region (R1) of default size was defined around the centre of the oocyst or cyst population (Figure 2). A rectangular sort region (R2) was defined around the respective bead standard used. Once sort regions were defined, a sample of water concentrate was analysed to allow the operator to increase or decrease the discriminator until extraneous particles collected fluoresced just below R1 (Figure 3). Once all regions were defined 5,000 events were collected for every sample. It was important that R1 be moved to the centre of the oocyst or cyst population for each antibody evaluated as fluorescence varies between mAbs.

#### Data Analysis

Data analysis was carried out using LYSIS II software obtained from Becton-Dickinson, (Lane Cove, NSW Australia). Bivariate dot-plots defining SSC versus FL1 were used for analyses (Figures 3 and 4). Regions were defined as above. To enumerate the number of beads detected, a sort rectangle (R2) was defined around the bead population. This region was determined by analysis of a control sample consisting of beads and oocysts/cysts (Figure 3). The specificity of each mAb was calculated in terms of a non-specific binding ratio by dividing the number of events collected in R1 by R2.

#### Statistical Analysis

Data interpretation was performed using Microsoft Excel™ 4.0 and the Analysis Toolpack™ Add-in. The hypothesis that the means from different samples were equal was tested using analysis of variance (ANOVA). Using a significance level of 5% ( $p < 0.05$ ), the critical values for the *F*-statistic were calculated and compared to that obtained from the ANOVA.

#### RESULTS

##### Analysis of mouse serum

Flow cytometric analysis of mouse serum revealed a large immunological response to the surface of *Cryptosporidium* oocysts in group E



mice (Table 1). Twenty-one days after the second immunisation both IgM and IgG *Cryptosporidium*-specific antibodies were detected in the serum of group E mice. Oocysts stained with serum from group E mice fluoresced very brightly even when the serum was diluted to 1 in 10.000 (Table 1). The fluorescence of oocysts stained with serum from group W mice before and 21 days after the second immunisation were not significantly ( $p > 0.05$ ) different indicating that the immunisation did not cause the production of antibodies specific to the surface of *Cryptosporidium* oocysts. No antibodies specific to the surface of *Cryptosporidium* oocysts were detected in the serum of group W mice. The slight difference between the results for group W and group E mice prior to immunisation was most probably due to variation in the sensitivity of the instrument on different days.

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**Table 1.** Comparison of the fluorescence intensity of *Cryptosporidium* oocysts stained with serum (diluted 1 in 1000) from group E and W mice and then stained with an anti-IgG or an anti-IgM fluorescently labelled antibody. Serum was tested prior to immunisation and then 21 days after the second immunisation.

mouse number	IgG specific antibody		IgM specific antibody	
	Pre-immunisation	Post-immunisation	Pre-immunisation	Post-immunisation
E1	12	1117	12	111
E2	9	460	11	121
E3	10	195	11	74
E4	10	1762	13	258
E5	12	550	12	33
W1	36	39	16	17
W2	45	32	27	14
W3	36	37	27	13
W4	42	30	25	27
W5	47	40	16	15
Positive* control (IgM)	42		978	
Negative Control	27		17	

ND - not determined

\* a IgM monoclonal antibody specific to *Cryptosporidium* was used as a positive control. A IgG positive control was not available.

### Hybridomas

Mouse number 4 from group E was sacrificed and the spleen cells fused with mouse myeloma cells. Screening of the resulting 230 clones identified six clones that were producing *Cryptosporidium*-specific antibody. Five of the clones (clones P9F1, P6G7, P6B11, P11D5 and P7D5) were found

to be producing IgM antibody specific to the surface of *Cryptosporidium* oocysts. One clone (CRY104) was producing IgG1 antibody specific to the surface of *Cryptosporidium* oocysts.

Additional fusions of spleen cells from mice in group E have produced a large number of hybridomas specific for the specific to the surface of *Cryptosporidium* oocysts. Further IgG1 antibodies produced by these hybridomas have been characterised. This further demonstrated that the methods according to the present invention are particularly suitable to produce useful IgG1 antibodies specific to the surface of *Cryptosporidium* oocysts.

Fusions were not attempted with mice from group W because of the poor immunological response to *Cryptosporidium* that was observed.

#### **Evaluation of monoclonal antibodies**

Results of flow cytometric analysis of water samples seeded with oocysts and stained with some of the *Cryptosporidium*-specific monoclonal antibodies are presented in Figure 1. Note the differences in the position of the population of oocysts along the Y axis. The oocyst population is closer to the top of the dotplot due for the sample stained with 8C12 (CRY104) than for any of the other samples. This is because the oocysts are fluorescing more brightly in this sample than any other sample. The fluorescence of the debris particles (below the oocysts) is not brighter in the 8C12 (CRY104) sample than in the unstained control. The separation between the oocysts and the debris particles is greatest in the 8C12 (CRY104) stained sample. This suggests that this antibody is most useful for staining *Cryptosporidium* oocysts in water samples. In comparison, the sample stained with 6G7 shows an increase in the fluorescence of some of the debris particles (to the right of the oocysts) when compared to the unstained control. This is due to this antibody binding to some of the debris particles and suggests that this antibody may not be useful for staining *Cryptosporidium* oocysts in water samples.

Flow cytometric analysis of serum from mice immunised with intact purified oocyst walls (group WI mice) revealed no difference in the brightness of *Cryptosporidium* oocysts stained with serum collected before or after immunisation (Table 2). Similar results were observed when using both IgM specific and IgG specific second antibodies. In comparison, when oocysts were stained with the serum from group WP mice there was a

difference in the fluorescence intensity of the oocysts stained with post immunisation serum and those stained with pre-immunisation serum. Results were similar for oocysts stained with both IgG specific and IgM specific secondary antibodies.

5

**Table 2.** Comparison of the fluorescence intensity of *Cryptosporidium* oocysts stained with serum (diluted 1 in 100 or 1 in 1000) from group WI and group WP mice and then stained with an anti-IgG or an anti-IgM fluorescently labelled antibody. Serum was tested prior to immunisation and then 21 days after the second immunisation.

10

mouse number	IgG specific antibody		IgM specific antibody	
	Pre-immunisation	Post-immunisation	Pre-immunisation	Post-immunisation
<sup>1</sup> WI1	36	39	16	17
WI2	45	32	27	14
WI3	36	37	27	13
WI4	42	30	25	27
WI5	47	40	16	15
<sup>2</sup> WP1	<sup>3</sup> 230	248	404	420
WP2	231	518	405	514
WP3	229	306	403	475
WP4	233	290	410	417
WP5	232	350	407	603

<sup>1</sup>WI - mice immunised with intact purified oocyst walls. Serum was diluted 1 in 1000.

<sup>2</sup>WP - mice immunised with small pieces of purified oocyst walls. Serum was diluted 1 in 100.

<sup>3</sup>Results of the two groups of mice are not directly comparable. Serum was not tested at the same dilution for both groups of mice. Different second antibodies were used to test the group W mice and the group WP mice.

15

### Antibody Comparison Tests

When concentrated water samples were stained with mAbs directed to either *Cryptosporidium* or *Giardia* there was a significant ( $p < 0.05$ ) increase in the number of fluorescent particles observed with the stained, compared with unstained samples (Table 5 and 6). Differences were observed between different antibodies with respect to the immunofluorescent staining, with increases in fluorescent particles detected between mAbs. The IgG1 antibody CRY104 resulted in significantly ( $p < 0.05$ ) lower levels of non-specific binding by the fluorescent mAb to extraneous particles in water, than when staining with antibodies of the IgM or IgG3 subclass (Table 5). The IgG1 antibody G203 produced specifically for water analysis of *Giardia*, resulted in significantly less fluorescent background particles than when staining with antibodies produced for faecal analysis including an IgG1 (Table 6).

Specific antibodies (both *Giardia* and *Cryptosporidium*) were selected for evaluation of the effect of mAbs on staining different water types. Statistically different levels ( $P < 0.05\%$ ) of non-specific binding of particles to antibodies were observed between different water types for both *Giardia* and *Cryptosporidium* analysis (Table 7 and 8). The variation observed following staining with an IgG3 was higher between waters than the variation observed by the IgG1 mAb (Table 7). Greater variation in non-specific binding was seen with Hydrofluor™ mAb compared with G203 (Table 8) thus significant differences in the level of binding between mAbs for every water type was observed.

**Table 3.** *Cryptosporidium*-specific monoclonal antibodies evaluated.

Antibody	Antibody Type	Supplier
CRY104	IgG1 FITC conjugate	MUCAB*
CRY26™	IgM FITC conjugate	MUCAB*
CRY212	IgM FITC conjugate	MUCAB*
Crypto-Cell	IgM FITC conjugate	CelLab, Australia
Crypto-a-glo™	IgM FITC conjugate	Waterborne. Inc. USA
Inmucell™	IgG3 FITC conjugate	Immucell, Portland. USA
Hydrofluor™	IgM Not conjugated	Meridian, ENSYS Inc. USA

\*Macquarie University Centre for Analytical Biotechnology

**Table 4.** *Giardia*-specific monoclonal antibodies evaluated.

Antibody	Antibody Type	Supplier
G203	IgG1 FITC conjugate	MUCAB*
Giardia-Cell	IgM FITC conjugate	CelLab, Australia
Giardia-a-glo™	IgG1 FITC conjugate	Waterborne. Inc. USA
Hydrofluor™	Giardia IgG Not	Meridian, ENSYS Inc.
Combo	conjugated + Crypto IgG	USA

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**Table 5.** Comparison of the level non-specific binding to *Cryptosporidium*-specific mAbs

Antibody	Non-specific Binding ratio*
CRY104	0.045 ± 0.013
CRY26	0.623 ± 0.023
CRY212	0.112 ± 0.034
Crypto-Cell	0.177 ± 0.034
Crypto-a-glo™	0.980 ± 0.125
Immucell™	0.155 ± 0.013
Hydrofluor™	0.159 ± 0.058
Unstained	0.033 ± 0.010

Results are the means of 3 separate analyses.

\* Arbitrary scale developed for *Cryptosporidium* analysis (see M&M)

Statistical difference (p<0.05) by ANOVA

**Table 6.** Comparison of the level of non-specific binding to *Giardia* specific mAbs

Antibody	Non-specific Binding ratio *
G203	0.029 ± 0.002
Giardia-Cell	0.163 ± 0.024
Giardia-a-glo™	0.201 ± 0.032
Hydrofluor™	0.044 ± 0.005
Unstained	0.020 ± 0.002

5 Results are the means of 3 separate analysis.

\* Arbitrary scale developed for *Giardia* Analysis (see M&M)

Statistical difference (p<0.05) by ANOVA

10 **Table 7.** Evaluation of the level of non-specific binding of *Cryptosporidium* specific mAbs in different water concentrates.

Water Type	Treatment mean ± SD		
	Unstained control	CRY104	Immucell™
Raw	0.023 ± 0.002	0.023 ± 0.001	3.294 ± 0.327
Backwash	0.023 ± 0.003	0.020 ± 0.006	0.440 ± 0.049
Raw	0.021 ± 0.003	0.020 ± 0.004	0.385 ± 0.016
Backwash	0.022 ± 0.002	0.087 ± 0.038	0.233 ± 0.052
Effluent	0.035 ± 0.006	0.085 ± 0.01	1.801 ± 0.093
Backwash	0.023 ± 0.004	0.277 ± 0.031	0.869 ± 0.154
Raw	0.022 ± 0.006	0.025 ± 0.005	1.391 ± 0.036
Raw	0.036 ± 0.006	0.099 ± 0.013	0.651 ± 0.047
Backwash	0.023 ± 0.004	0.031 ± 0.002	0.156 ± 0.027
Filtered	0.026 ± 0.007	0.027 ± 0.002	0.166 ± 0.015

Statistical difference (p<0.05) by ANOVA, N=3

**Table 8.** Evaluation of the level of non-specific binding of *Giardia* specific mAbs in different water concentrates.

Water Type	Treatment mean $\pm$ SD		
	Unstained control	G203	Hydrofluor™
Raw	0.022 $\pm$ 0.002	0.048 $\pm$ 0.007	0.645 $\pm$ 0.01
Backwash	0.022 $\pm$ 0.003	0.034 $\pm$ 0.005	0.669 $\pm$ 0.008
Raw	0.020 $\pm$ 0.003	0.025 $\pm$ 0.002	0.696 $\pm$ 0.022
Backwash	0.021 $\pm$ 0.003	0.214 $\pm$ 0.094	14.63 $\pm$ 4.58
Effluent	0.032 $\pm$ 0.007	0.007 $\pm$ 0.001	1.021 $\pm$ 0.075
Backwash	0.021 $\pm$ 0.003	0.081 $\pm$ 0.014	10.36 $\pm$ 1.547
Raw	0.021 $\pm$ 0.006	0.023 $\pm$ 0.003	0.062 $\pm$ 0.005
Raw	0.031 $\pm$ 0.006	0.070 $\pm$ 0.022	1.637 $\pm$ 0.096
Backwash	0.022 $\pm$ 0.004	0.036 $\pm$ 0.009	0.111 $\pm$ 0.016
Filtered	0.024 $\pm$ 0.006	0.024 $\pm$ 0.003	0.043 $\pm$ 0.004

5 Statistical difference ( $p < 0.05$ ) by ANOVA, N=3

Evaluation of the number of fluorescent particles within a water concentrate before and after immunofluorescent staining revealed that with all mAbs tested there was significantly ( $p < 0.05$ ) higher numbers of fluorescent particles following staining. There was, however, a large variation in the number of fluorescent particles after staining with different mAbs. Analysis of different classes of mAbs revealed that staining with the IgG1 mAbs generally resulted in less fluorescent particles than staining with IgM and IgG3 mAbs.

15 Variation in the background fluorescence observed is largely due to the type of antibody used for the assay. The primary immune response to an antigen induces the production of IgM. Immunoglobulin M antibodies exist as large mAbs containing ten binding sites, with a greater chance of binding non-specifically to particles compared with IgG. The surface of oocysts and cysts contain carbohydrates that induce the production of low affinity IgM and IgG3. Immunoglobulin G1 mAbs are generally produced following repeated exposure to an antigen. Isotype switching occurs from IgM to IgG, resulting in higher affinity mAbs containing two identical binding sites more



specific to the antigen they are produced to. Specificity of the IgM, CRY212 that was developed for water analysis was greater than commercial mAbs that were produced for clinical applications such as faecal analysis. This trend was observed following *Giardia* analysis of IgG1 mAbs, with G203 having higher affinity than Giardia-a-glo™.

Comparison of different water types revealed a significant difference in the non-specific binding ratio between waters. Variation was observed following staining with IgG1 mAbs (CRY104 and G203). however, the range of non-specific binding to different water types was significantly reduced over the other mAbs examined. The degree of variation in non-specific binding was significant with backwash samples displaying a diverse range in background fluorescence between a ratio of 0.111 and 14.63 for non-specific binding of the Hydroflor mAb. Thus, some water concentrates contain particles that have a much higher chance of binding to certain mAbs, however, for all water types the IgG1 mAbs caused the least increase in the number of fluorescent particles detected.

mAbs produced specifically for water analysis have a high affinity for cysts and oocysts over extraneous particles present within water concentrates. Immunoglobulin G1 are smaller mAbs that bind specifically to antigenic sites that they are produced to with little background interference. Production of new IgG1 mAbs to different antigenic sites on cysts or oocysts may enable the development of highly specific assays for *Cryptosporidium* and *Giardia* detection within environmental samples.

**Table 9.** Comparison of the fluorescence intensity of *Cryptosporidium* oocysts stained with serum of three dilutions (1:1000, 1:10,000, 1:100,000) from the mouse groups E (Extract), W (Walls) oocyst control (OC - mice injected with whole oocysts) or negative control (C) and then stained with an anti-IgG or an anti-IgM fluorescently labelled antibody. Data was calculated by subtracting the fluorescent value (arb) obtained from second serum samples after two immunisations from the negative serum samples prior to immunisation.

	IgM Specific Antibody			IgG Specific Antibody		
	1:1000	1:10,000	1:100,000	1:1000	1:10,000	1:100,000
Extract						
Mouse 1	280	129	64	353	370	229
Extract						
Mouse 2	283	115	96	220	259	172
Extract						73
Mouse 3	278	129	90	67	84	
Extract						
Mouse 4	319	151	108	354	425	279
Extract						
Mouse 5	45	92	99	103	324	44
Walls						
Mouse 1	3	30	19	-68	-13	28
Walls						
Mouse 2	-3	36	2	29	-31	10
Walls						
Mouse 3	18	0	4	41	52	16
Walls						
Mouse 4	38	38	-3	24	-3	-4
Walls						
mouse 5	22	16	15	3	-11	27
Oocyst						
Control 1	432	240	115	2	-18	13
Oocyst						
Control 2	339	189	26	-53	-5	42
Oocyst						
Control 3	438	228	69	48	45	2
Oocyst						
Control 4	581	457	235	124	46	57
Oocyst						
Control 5	210	75	37	21	43	43
Control	0	0	0	0	0	0

### Western Blot Examination of serum samples and mAbs

A large amount of immunostained bands were observed in western blots carried out on mouse serum from each of the three immunisation groups (E, W, and OC). The oocyst control mouse group (OC) and extract mouse group (E) showed banding from > 200 kD down to 40kD with no distinct difference in banding patterns. The wall mouse group (W) showed unique banding only in the mid range between 100 kD and 160 kD. mAbs analysed by western blot showed some unique banding patterns although all recognised two distinct bands, one at 200 kD and the other at 40 kD.

### Functional Measurement of Avidity

**Table 10:** Shows the functional measurement of Avidity.

Monoclonal Antibody	Subclass	ka (affinity constant of whole antibody)
CRY104	IgG1	$7 \times 10^8 \text{ mol}^{-1}$
CRY26	IgM	$3 \times 10^8 \text{ mol}^{-1}$
Immucell™	IgG3	$6 \times 10^7 \text{ mol}^{-1}$

During the course of the immunisation program mice were monitored for both IgM and IgG levels in serum. The oocyst control group demonstrated a high IgM response with little or no IgG. This would suggest that there is no memory in the immune response (i.e. shift to IgG response) and that each immunisation is seen as a new antigen.

This may be due to *C. parvum* oocysts containing complex polysaccharide structures which do not require help of T-Lymphocytes to stimulate B-Cells into antibody production. Thus resulting in less isotype switching and reduced affinity maturation in mAbs produced. Hence less specific IgM antibodies dominate the immune response.

The oocyst walls group (W) showed no immune response. This is probably due to the large size and structure of the whole oocyst wall which may have a desensitising effect on the immune response. The fact that no immune response was seen in the oocyst wall group would suggest that with no sporozoites present a decrease in immunogenicity is obtained.

The oocyst extract group elicited strong IgM and IgG responses. IgM responses, however, were not as strong as in the oocyst control group, again

suggesting that sporozoites are more antigenic than the oocyst wall and are characteristic of the self limiting nature of *Cryptosporidiosis*. It appears that the SDS digestion breaks up the oocyst wall and exposes more epitopes for an immune response. After removing the sporozoites and breaking up the oocyst wall structure into smaller complexes the oocyst wall or extract becomes an extremely immunogenic antigen as shown from the strong immune response seen in the extract mice. After further immunisations in extract mice the IgG response further increased as for a typical immunisation program. This was not observed in the Control or Wall group, which showed dominant IgM response or no response respectively.

From the western blot analysis, extract and oocyst control groups share a similar banding pattern, with extract mice showing a more definitive banding pattern with one lower distinct band unshared (40 kD).

Measurement of the antibody avidity was made and compared to other anti-Crypto antibodies. The affinity constant for the IgG1 whole antibody was calculated to be  $7 \times 10^8 \text{ mol}^{-1}$ , which is at least double that of the other antibodies tested (Table 10). This indicates that CRY104 is at least twice as effective at binding to *Cryptosporidium* at a constant concentration than the other antibodies tested, and forms a stronger bond to the epitope. It was found that the binding efficiency of CRY104 facilitates more effective staining of *Cryptosporidium* at lower concentrations than other commonly used antibodies.

#### CONCLUSIONS

A strong immunological response to the surface of *Cryptosporidium* oocysts was produced by immunising mice with a partially purified sample of the outer layer of the oocyst wall. Immunising mice with purified oocyst walls did not produce a strong immunological response. This would suggest that there is either immune suppression by a component of the oocyst wall or that the natural presentation of the surface antigens on the oocyst walls does not produce an immunological response. The present inventors have overcome this lack of an immunological response to the surface of *Cryptosporidium* oocysts by partially purifying the surface antigen.

Fusion of spleen cells from one mouse immunised with the purified sample of the outer layer of the oocyst wall resulted in six (6) hybridomas that produce antibody that is specific to the surface of the oocyst wall. One of these six antibodies is of the IgG1 immunological subclass, the remaining

five antibodies are of the IgM subclass. The IgG1 antibody appears to be superior to the IgM antibodies for staining *Cryptosporidium* oocysts in water samples. Mice immunised with purified *Cryptosporidium* oocysts walls do not produce a strong immunological response to the surface of *Cryptosporidium* oocysts. If, however, the purified oocysts walls are broken up into small pieces, then a strong response is produced in both the IgG and IgM immunological subclasses. Mice immunised with such a procedure would also be suitable for producing monoclonal antibodies of the IgG1 subclasses.

It will be appreciated by persons skilled in the art that numerous variations and/or modifications may be made to the invention as shown in the specific embodiments without departing from the spirit or scope of the invention as broadly described. The present embodiments are, therefore, to be considered in all respects as illustrative and not restrictive.